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This listing of the claims will replace all prior versions and listings of claims in the application:

## Listing of the claims:

Claim 1: (currently amended) A method for identifying an agent er event—capable of priming a cell for preconditioning and/or inducing preconditioning of a cell, tissue or organ comprising assessing the ability of the agent er event—to modulate abundance of a preconditioning protein in a cell, tissue or organ by detecting a modulation in abundance of the preconditioning protein in the presence of the agent er event—as compared to the abundance of preconditioning protein in the absence of the agent—er—event, wherein the preconditioning protein is a protein of an oxidative phosphorylation (OxPhos) pathway, tricarboxylic acid (TCA) cycle, a Ca<sup>2+</sup> handling protein, a chaperone protein, or a protein selected from aldehyde dehydrogenase, NG-dimethylarginine dimethylaminohydrolase (DDAH), and the RNA binding protein regulatory subunit DJ-1.

Claim 2-12: (canceled)

Claim 13: (currently amended) The method of claim 1 wherein the agent or event-identified modulates the

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abundance of preconditioning protein in the cell, tissue or organ and leads to a change via cross-talking, a feed-back mechanism or a signaling mechanism which effects the first window of preconditioning, the second window of preconditioning or both windows of preconditioning of a cell.

Claim 14: (currently amended) The method of claim 1 wherein the agent or event identified modulates the abundance of preconditioning protein in the cell, tissue or organ and leads to a change in function of a protein complex or pathway of which the modified protein is a member.

Claim 15: (currently amended) The method of claim 1 wherein the agent or event identified modifies a mitochondrial protein.

Claim 16: (currently amended) The method of claim 1 wherein the agent or event identified increases a level the abundance of one or more of isocitrate dehydrogenase NAD+\_ specific subunit alpha IDH, succinyl CoA ligase, a 23 kDa mitochondrial precursor subunit of Complex I, a 24 kDa mitochondrial precursor subunit of Complex I, a 30 kDa

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mitochondrial precursor subunit of Complex I, a  $\delta$  chain mitochondrial precursor of an F1 portion, a d chain mitochondrial precursor of a Fo portion of Complex V, prohibitin, ADP ribosyl hydrolase, HSP27 and RNA binding protein regulatory subunit (DJ-1).

Claim 17: (currently amended) The method of claim 1 wherein the agent er event identified decreases a level the abundance of one or more of dihydrolipoamide succinyltransferase, core protein I of Complex III, metaxin 2 and sarcalumenin.

Claim 18: (currently amended) The method of claim 1 wherein the agent or event identified changes a level the abundance of DDAH.

Claim 19: (currently amended) The method of claim 1 wherein the agent or event identified increases posttranslational modification of  $\beta$  chain mitochondrial precursor of the F1 portion of Complex V, protein X, or aconitate hydratase (aconitase).

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Claim 20: (currently amended) The method of claim 1 wherein the agent or event mimics modulation of the preconditioning proteins by adenosine or diazoxide.